In the Claims

Applicant has submitted a new complete claim set, with insertions and deletions in amended claims indicated by underlining and strikeouts, respectively.

- 1. (Currently amended) A method for <u>identifying designing</u> a compound capable of binding to an active site, an accessory binding site or a pocket of an <u>ribosome recycling factor (RRF)</u> protein, which comprises computationally evaluating a chemical entity of RRF protein on the basis of a structure coordinate obtained from an RRF protein crystal <u>comprising steps of:</u>
- a) using a three-dimensional structure of said protein as defined by atomic coordinates of RRF protein according to Table 8;
- b) employing said three-dimensional structure to design or select said compound capable of binding to RRF protein;
 - c) synthesizing said compound capable of binding to RRF protein; and
- d) contacting said compound capable of binding to RRF protein with said RRF protein in the presence of a substrate to determine the ability of said compound capable of binding to RRF protein to bind said RRF protein.
- 2.-51. (Canceled)
- 52. (New) The method according to claim 1, wherein said compound capable of binding to RRF protein is designed de novo.
- 53. (New) The method according to claim 1, wherein said compound capable of binding to RRF protein is designed from a known compound capable of binding to RRF protein.